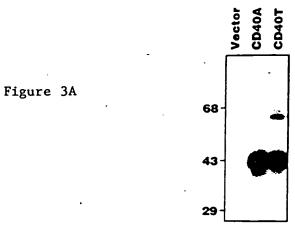
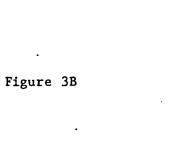
CD40T
CD40A
TNFR
FAS
ATNFR
E12
SNF1

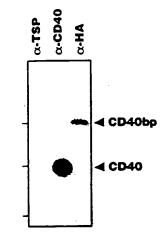
Figure 2

Figure 1

	In-vitro- translated product		GST		GST CD40T		GST CD40A	
, .	Luciferace	CD40 BD	Luciferace	CD40 BD	Luciferace	CD40 BD	Luciferace	CD40 BD
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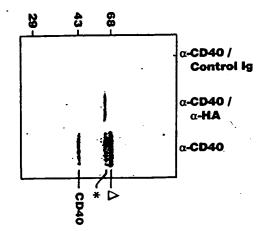
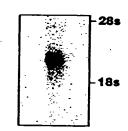


Figure 3C

Figure 3D



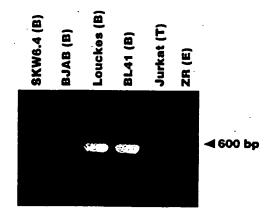


Figure 3E

165 220 275 330 385 54 495 TACQESIVKDKVFKDNÇÇKREI ELPÇVRPDÇKEKVLRKDLRDHV D # V S A # SLY 71 X1 0 FKKPTGEMNIASGCPVFVAQTVLENGTYIKD 의 지 O 의 기 기 FVIMRGEYDALLPWPFKQKVT LRINNESKILHLORVIDSOAE ESVDKSAGOVARNTGLES WKIRDYKRRKOEAVMGKTT KTL EL **太| 田| の** ഗ SVQTLLR PÇVVVSÇPHKÇS ASSAVQHVNLLK FVP ELEROKEMUR SLONRVTELE TASYNGVLIE PKQTECGHRFCESCMAALLSSSPKCTRNESRGÇAEQLMLGHLVHLKNDÇHFEIEATÇSHÇKSQVPMIALQKHEDTDÇPÇ KGTHLSLF <u>></u> SAGT मा वा व S TDR S 国工工 GYKMCARVYLNGDG ഗ HINOUR CORRESPONDENCE OF CORRE DPNS NPPLKLH SRRHLGDAFKP DLPD PGALQT TIFIKVIVDTS SSKKMDS >-CHLVLCSE LALQIYÇE 기 H J 의 도 S OPFYTG MLMDQGS œ. EÇVNA EKAÇKY (의 조) (이 (의 Σ ZIOIO L S 기 이 기 포 回

Figure 4A

figure 4B

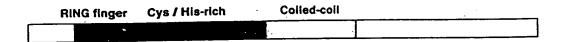
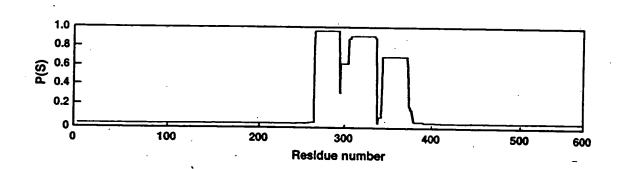


Figure 4D



CD40bp TRAF2 CD40bp TRAF2 CD40bp TRAF2 TRAF1 TRAF2 TRAF1 TRAF2 **TRAF1 TRAF1** ن ہے بد _ > < , Z I يعتعدا $x \propto x$ ۵ ، ی **500** v zz FITS FXXIS FISHTS م م م v ziz ××× ××× ×××× بنى ۳ که ۲ ۲ ح > -TASYNGVE VSTYDGVE EASIF DGTF 0.00 4 777 8 8 8 2002 244 244 244 244 200 A E Y I G Y F G Y K M A E Y I S R Y G Y K M A F Y T A K Y G Y K L 000 000 000 444 ×z× யீழ் HH> ZZZ XX+ o o o r min م 0 **E E E** A L L P ш С ш o เกิเท -> ° しるま ு ய ய م به ت 000 $\times \propto \propto$ > z > w ₀ w SON ×00 >>> $\circ \circ \times$ **≻ 14 14** ii.u.u. **4 4** 0 ئز ب ب ပ ပ ပ SON NAN $\alpha \times \alpha$ SOA SSS **-** 0 $\vdash\vdash\vdash$ H > H ے تے ہ × & & ェネネ ی ی ی 000 نا با با گ ے ہے۔ Z Q U SOA × & & х қ х о о о >>> 4 4 V SSS 4 4 4 0 0 0 ن ب ب 0 × 0 OOI xxx 326 234 406 366 509 445 484 547

Figure .5

CACTIGGTTAAGGTCCCAGAGCTCAGAATCAGAATCAGAATCAGATCAGAAACCTGGCTCCTGGCTCCTGCTCCTGCTCCTTCTTAAGG atcgctistcctisacagaaagaacticctctttcctaaaatggagticgagtaaaaagagaacticticctgggggggg CCGCCGCTAAAGCTGCACACTGACCGCAGTGCTGGGACCCCAGTTTTTTGTCCCTGAACAAGGAGGTTACAAGGAAAAGTTTGTGAA **ACAPAAGGAAATGCTTCGAAATAATGAATCCAAAATCCTTCATTTACAGCGAGTGATAGACAGCCAAGCAGAGAAACTGAAGGAG** PTGACAAGGAGGGCGCTTTCCGGCAGAACTGGGAAAGGAAGCAGACAGCAGCATGAAGAGCAGCGTGGAGTCCCTCCAGAAACGGGTG :CCGAGCTGGAGAGCGTGGACAAGAGCGCCGGGCAAGTGGCTCGGAACACAGGCCTGCTGGAGTCCCAGCTGAGCCGGCATGACCA <u> BATIOCTIGAGTIGTIGCACGACATICCGCCTAGCCGACATIGGACCTTCCAGGTICCTGGAGACCGCCAGCTACAATIGGAGTIGCTICA</u> ITCTTCCTTAAACTTGAACAÇCAAAAAAACACACACACACACACACGTGGGGATAGCTGGACATGTCAGCATGTTAAGTAAAAGA GACCCTGGAGGACAAGTACAAGTGTGAGAAGTGCCACCTGGTGCTGTGCAGCCCGAAGCAGACCGAGTGTGTGGGCACCTTCTGCG SATAATTGCTGCAAGAGAGAÄTTCTGGCTCTTCAGATCTATTGTCGGAATGAAÄGCAGAGGTTGTGCAGAGCAGTTAATGCTGG <u> ACCTGCGAGACCACCTGGAGAAGGCGTGTAAATACCGGGAAGCCACATGCAGCCACTGCAAGAGAGTCAGGTTCCGATGATGATCGCGCTG</u> <u> BAGTGCACACTTGTCAGAGTGTGTGAATGCCCCCAGCACCTGTAGTTTTTAAGCGCTATGCGCTGCGTTTTTTCAGGGGACAAACCAGC</u> ITGTTGCAGAATGAAAGTGTAGAAAAAAAAACAAGACATACAAAGTTTGCACAATCAGATATGTAGCTTTGAAAATTGAAATTGAGAG SCTTACITTGCCTATAAGATGTGTCCCAGGGTCTACCTGAACGGGGACGGGATGGGGGAAGGGGACGCACTTGTCGCTGTTTTTTTGT ITICGGAICTGCCCCAITCCCTGATAAGTAGCTGGGAGGTGGATTTAACAGAAAGGCAACTCCTCTGGGGGATTTTGAACCGGTCTGTC IGICAGAGAAGGTITITCATITITCATTTTAAGATCTAGTTAATTAAGGTGGAAAACATATATGCTAAACAAAAGAAACATGATTT <u> ACGANGGCCACGCCCCGGCCCCTGAGCCGGCCGAGCGGCGACGGACCGCGAGATGAGGAAPATGAGGCCCAAAGAAGTGATGATGC</u> CAGAAACACGAAGAACACCGACTGTCCCTGCGTGGTGGTCCTGCCCTCACAAGTGCAGGCGTCCAGACTCTCCTGAGGAGGGGAGTT CATCATGCGTGGAGAATATGATGCCCTGCTTCCTTGGCCGTTTAAGCAGAAAGTGACACTCATGCTGATGGATCAGGGGGTCCTCTC **Baatitatgaaatagta**